IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In the Application of:

OMOLAYO O. FAMODU ET AL.

CASE NO.: BB1395 US PCT

APPLICATION NO.: 10/069,427

GROUP ART UNIT: UNKNOWN

FILED: UNKNOWN

EXAMINER: UNKNOWN

INTERNATIONAL APPLICATION NO.: PCT/US00/26442

INTERNATIONAL FILING DATE: SEPTEMBER 9, 2000

FOR: GENES ENCODING STEROL

DELTA-15 REDUCTASE IN PLANTS

STATEMENT UNDER 37 CFR 1.821(g) and 1.825(b)

Commissioner for Patents, Box PCT United States Patent and Trademark Office Washington, DC 20231

Sir:

The submission of the substitute Sequence Listing filed concurrently herewith does not include new matter.

The copy of the substitute Sequence Listing in computer readable form filed concurrently herewith is the same as the paper copy of the substitute Sequence Listing filed concurrently herewith.

Respectfully submitted,

Nou & Slauder Lori Y Beardell Attorney D Attorney For Applicants Registration No. 34,293

Telephone: 302-992-4926 Facsimile: 302-892-1026

Dated: July 11, 2002

10/069427

SEQUENCE LISTING

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<110> Famodu, Omolayo O.
       Kinney, Anthony J.
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 <130> BB1395 PCT
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Val Pro Gly Val Ala Leu Leu Asp Gly Thr Arg Leu His Tyr Cys Cys
Asn Gly Leu Leu Ser Leu Leu Leu Val Ala Leu Leu Gly Ile Gly
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Ala Lys Met Gly Phe Val Ser Pro Thr Ala Ile Ser Asn Arg Gly Leu
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Glu Leu Leu Ser Thr Thr Phe Ala Phe Ser Phe Leu Val Thr Leu Ile 85 90 95

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- Leu Leu Asp Gly Thr Arg Leu His Tyr Cys Cys Asn Gly Leu Leu Ser 50 55 60
- Leu Leu Leu Val Ala Leu Leu Gly Ile Gly Ala Lys Met Gly Phe 65 70 75 80
- Val Ser Pro Thr Ala Ile Ser Asp Arg Gly Leu Glu Leu Leu Ser Thr 85 90 95
- Thr Phe Ala Phe Ser Phe Leu Val Thr Leu Ile Leu His Phe Ser Gly 100 105 110
- Cys Lys Ser Gln Ser Lys Gly Ser Ser Leu Lys Pro His Leu Ser Gly 115 120 125
- Asn Leu Ile His Asp Trp Trp Phe Gly Ile Gln Leu Asn Pro Gln Phe 130 135 140
- Met Gly Ile Asp Leu Lys Phe Phe Phe Val Arg Ala Gly Met Met Gly 145 150 155 160
- Trp Leu Leu Ile Asn Leu Ser Ile Leu Met Lys Ser Ile Gln Asp Gly
 165 170 175
- Thr Leu Ser Gln Ser Met Ile Leu Tyr Gln Leu Phe Cys Ala Leu Tyr 180 185 190
- Ile Leu Asp Tyr Phe Val His Glu Glu Tyr Met Thr Ser Thr Trp Asp 195 200 205
- Ile Ile Ala Glu Arg Leu Gly Phe Met Leu Val Phe Gly Asp Leu Val 210 215 220
- Trp Ile Pro Phe Ser Phe Ser Ile Gln Gly Trp Trp Leu Leu Met Asn 225 230 235 240
- Ser Val Glu Leu Thr Pro Ala Ala Ile Val Ala Asn Cys Phe Val Phe 245 250 255
- Leu Ile Gly Tyr Met Val Phe Arg Gly Ala Asn Lys Gln Lys His Val 260 265 270
- Phe Lys Lys Asn Pro Lys Ala Pro Ile Trp Gly Lys Pro Pro Lys Val 275 280 285
- Ile Gly Gly Lys Leu Leu Ala Ser Gly Tyr Trp Gly Ile Ala Arg His 290 295 300
- Cys Asn Tyr Leu Gly Asp Leu Met Leu Ala Leu Ser Phe Ser Leu Pro 305 310 315 320
- Cys Gly Ile Ser Ser Pro Ile Pro Tyr Phe Tyr Pro Ile Tyr Leu Leu 325 330 335
- Ile Leu Leu Ile Trp Arg Glu Arg Arg Asp Glu Ala Arg Cys Ala Glu 340 345 350

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Gly Lys Pro Pro Lys Val Val Gly Gly Lys Leu Leu Ala Ser Gly Tyr 50 60

Trp Gly Ile Ala Arg His Cys Asn Tyr Leu Gly Asp Leu Leu Ala 65 70 75 80

Leu Ser Phe Ser Leu Pro Cys Gly Val Ser Ser Val Val Pro Tyr Phe 85 90 95

Tyr Pro Thr Tyr Leu Leu Ile Leu Leu Val Leu Arg Glu Arg Arg Asp 100 105 110

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Met Ile Leu Tyr Gln Leu Phe Cys Ala Leu Tyr Ile Leu Asp Tyr Phe 180 185 190

Val His Glu Glu Tyr Met Thr Ser Thr Trp Asp Ile Ile Ala Glu Arg 195 200 205

Leu Gly Phe Met Leu Val Phe Gly Asp Leu Val Trp Ile Pro Phe Ser 210 225 220

Phe Ser Ile Gln Gly Trp Trp Leu Leu Met Asn Ser Val Glu Leu Thr 225 230 235

Pro Ala Ala Ile Val Ala Asn Cys Phe Val Phe Leu Ile Gly Tyr Met 245 250 255

Val Phe Arg Gly Ala Asn Lys Gln Lys His Val Phe Lys Lys Asn Pro 260 265 270

Lys Ala Pro Ile Trp Gly Lys Pro Pro Lys Val Ile Gly Gly Lys Leu 275 280 285

Leu Ala Ser Gly Tyr Trp Gly Ile Ala Arg His Cys Asn Tyr Leu Gly 290 295 300

Asp Leu Met Leu Ala Leu Ser Phe Ser Leu Pro Cys Gly Ile Ser Ser 305 310 315 320

Pro Ile Pro Tyr Phe Tyr Pro Ile Tyr Leu Leu Ile Leu Leu Ile Trp 325 330 335

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His Leu Lys Phe Met Thr Thr Lys Ser Gly Ser Lys Leu Leu Ile Ser

340 345 350

Gly Trp Trp Gly Val Ala Arg His Val Asn Tyr Phe Gly Asp Trp Ile 355 360 365

Met Ala Trp Ser Tyr Cys Leu Thr Thr Gly Phe Asn Thr Pro Leu Thr 370° 375 380

Tyr Phe Tyr Val Ile Tyr Phe Gly Ile Leu Leu His Arg Asp Arg 385 390 395 400

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Ile Cys Ala Lys Leu Gly Ile Val Ser Pro Leu Val Val Ala Asp Arg 65 70 75 80

Gly Leu Glu Leu Ser Ala Thr Phe Ile Phe Cys Val Leu Val Thr 85 90 95

Leu Ala Leu Tyr Val Thr Gly Arg Ser Ser Ser Asn Lys Gly Ser Ser 100 105 110

Leu Lys Pro His Val Ser Gly Asn Leu Val His Asp Trp Trp Phe Gly 115 120 125

Ile Gln Leu Asn Pro Gln Phe Met Ser Ile Asp Leu Lys Phe Phe 130 135 140

Val Arg Ala Gly Met Met Gly Trp Leu Leu Ile Asn Leu Ser Ile Leu 145 150 155 160

Ala Lys Ser Val Gln Asp Gly Ser Leu Ser Gln Ser Met Ile Leu Tyr 165 170 175

Gln Ile Phe Cys Ala Leu Tyr Ile Leu Asp Tyr Phe Val His Glu Glu 180 185 190

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